

SEQUENCE LISTING

<110> Weiner, Howard L.  
Maron, Ruth  
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<120> SUPPRESSION OF VASCULAR DISORDERS BY  
MUCOSAL ADMINISTRATION OF HEAT SHOCK PROTEIN PEPTIDES

<130> B0801/7202 (AWS)

<150> US 60/189,855

<151> 2000-03-15

<160> 3

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 540

<212> PRT

<213> M. tuberculosis

<400> 1

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Lys	Gly	Arg	Asn	Val	Val	Leu	Glu	Lys	Lys	Trp	Gly	Ala	Pro	Thr	Ile
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Thr	Asn	Asp	Gly	Val	Ser	Ile	Ala	Lys	Glu	Ile	Glu	Leu	Glu	Asp	Pro
	50					55					60				
Tyr	Glu	Lys	Ile	Gly	Ala	Glu	Leu	Val	Lys	Glu	Val	Ala	Lys	Lys	Thr
65					70					75					80
Asp	Asp	Val	Ala	Gly	Asp	Gly	Thr	Thr	Thr	Ala	Thr	Val	Leu	Ala	Gln
				85					90					95	
Ala	Leu	Val	Arg	Glu	Gly	Leu	Arg	Asn	Val	Ala	Ala	Gly	Ala	Asn	Pro
			100					105					110		
Leu	Gly	Leu	Lys	Arg	Gly	Ile	Glu	Lys	Ala	Val	Glu	Lys	Val	Thr	Glu
		115					120					125			
Thr	Leu	Leu	Lys	Gly	Ala	Lys	Glu	Val	Glu	Thr	Lys	Glu	Gln	Ile	Ala
	130					135					140				
Ala	Thr	Ala	Ala	Ile	Ser	Ala	Gly	Asp	Gln	Ser	Ile	Gly	Asp	Leu	Ile
145					150					155					160
Ala	Glu	Ala	Met	Asp	Lys	Val	Gly	Asn	Glu	Gly	Val	Ile	Thr	Val	Glu
			165					170						175	
Glu	Ser	Asn	Thr	Phe	Gly	Leu	Gln	Leu	Glu	Leu	Thr	Glu	Gly	Met	Arg
		180					185						190		
Phe	Asp	Lys	Gly	Tyr	Ile	Ser	Gly	Tyr	Phe	Val	Thr	Asp	Pro	Glu	Arg
	195						200					205			
Gln	Glu	Ala	Val	Leu	Glu	Asp	Pro	Tyr	Ile	Leu	Leu	Val	Ser	Ser	Lys
	210					215					220				
Val	Ser	Thr	Val	Lys	Asp	Leu	Leu	Pro	Leu	Leu	Glu	Lys	Val	Ile	Gly
225					230					235					240
Ala	Gly	Lys	Pro	Leu	Leu	Ile	Ile	Ala	Glu	Asp	Val	Glu	Gly	Glu	Ala
			245						250					255	
Leu	Ser	Thr	Leu	Val	Val	Asn	Lys	Ile	Arg	Gly	Thr	Phe	Lys	Ser	Val
			260					265					270		



145					150					155				160
Pro	Val	Thr	Thr	Pro	Glu	Glu	Ile	Ala	Gln	Val	Ala	Thr	Ile	Ser
				165					170					175
Asn	Gly	Asp	Lys	Glu	Ile	Gly	Asn	Ile	Ser	Asp	Ala	Met	Lys	Lys
			180					185					190	
Val	Gly	Arg	Lys	Gly	Val	Ile	Thr	Val	Lys	Asp	Gly	Lys	Thr	Leu
		195					200					205		Asn
Asp	Glu	Leu	Glu	Ile	Ile	Glu	Gly	Met	Lys	Phe	Asp	Arg	Gly	Tyr
	210					215					220			Ile
Ser	Pro	Tyr	Phe	Ile	Asn	Thr	Ser	Lys	Gly	Gln	Lys	Cys	Glu	Phe
	225				230					235				Gln
Asp	Ala	Tyr	Val	Leu	Leu	Ser	Glu	Lys	Lys	Ile	Ser	Ser	Ile	Gln
			245						250					255
Ile	Val	Pro	Ala	Leu	Glu	Ile	Ala	Asn	Ala	His	Arg	Lys	Pro	Leu
			260					265					270	Val
Ile	Ile	Ala	Glu	Asp	Val	Asp	Gly	Glu	Ala	Leu	Ser	Thr	Leu	Val
		275					280					285		Leu
Asn	Arg	Leu	Lys	Val	Gly	Leu	Gln	Val	Val	Ala	Val	Lys	Ala	Pro
	290				295						300			Gly
Phe	Gly	Asp	Asn	Arg	Lys	Asn	Gln	Leu	Lys	Asp	Met	Ala	Ile	Ala
	305				310					315				320
Gly	Gly	Ala	Val	Phe	Gly	Glu	Glu	Gly	Leu	Thr	Leu	Asn	Leu	Glu
			325						330					335
Val	Gln	Pro	His	Asp	Leu	Gly	Lys	Val	Gly	Glu	Val	Ile	Val	Thr
			340					345					350	Lys
Asp	Asp	Ala	Met	Leu	Leu	Lys	Gly	Lys	Gly	Asp	Lys	Ala	Gln	Ile
		355					360					365		Glu
Lys	Arg	Ile	Gln	Glu	Ile	Ile	Glu	Gln	Leu	Asp	Val	Thr	Thr	Ser
	370					375					380			Glu
Tyr	Glu	Lys	Glu	Lys	Leu	Asn	Glu	Arg	Leu	Ala	Lys	Leu	Ser	Asp
	385				390					395				Gly
Val	Ala	Val	Leu	Lys	Val	Gly	Gly	Thr	Ser	Asp	Val	Glu	Val	Asn
			405						410					415
Lys	Lys	Asp	Arg	Val	Thr	Asp	Ala	Leu	Asn	Ala	Thr	Arg	Ala	Ala
		420					425						430	Val
Glu	Glu	Gly	Ile	Val	Leu	Gly	Gly	Cys	Ala	Leu	Leu	Arg	Cys	Ile
		435					440					445		
Pro	Ala	Leu	Asp	Ser	Leu	Thr	Pro	Ala	Asn	Glu	Asp	Gln	Lys	Ile
	450					455					460			Gly
Ile	Glu	Ile	Ile	Lys	Arg	Thr	Leu	Lys	Ile	Pro	Ala	Met	Thr	Ile
	465				470					475				480
Lys	Asn	Ala	Gly	Val	Glu	Gly	Ser	Leu	Ile	Val	Glu	Lys	Ile	Met
			485						490					495
Ser	Ser	Ser	Glu	Val	Gly	Tyr	Asp	Ala	Met	Ala	Gly	Asp	Phe	Val
			500					505					510	Asn
Met	Val	Glu	Lys	Gly	Ile	Ile	Asp	Pro	Thr	Lys	Val	Val	Arg	Thr
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Leu	Leu	Asp	Ala	Ala	Gly	Val	Ala	Ser	Leu	Leu	Thr	Thr	Ala	Glu
	530					535						540		Val
Val	Val	Thr	Glu	Ile	Pro	Lys	Glu	Glu	Lys	Asp	Pro	Gly	Met	Gly
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Met	Gly	Gly	Met	Gly	Gly	Gly	Met	Gly	Gly	Gly	Met	Phe		
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<210> 3  
 <211> 544  
 <212> PRT  
 <213> C. pneumoniae

<400> 3

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			20					25					30		
Pro	Lys	Gly	Arg	His	Val	Val	Ile	Asp	Lys	Ser	Phe	Gly	Ser	Pro	Gln
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Val	Thr	Lys	Asp	Gly	Val	Thr	Val	Ala	Lys	Glu	Ile	Glu	Leu	Glu	Asp
	50					55					60				
Lys	His	Glu	Asn	Met	Gly	Ala	Gln	Met	Val	Lys	Glu	Val	Ala	Ser	Lys
65				70					75					80	
Thr	Ala	Asp	Lys	Ala	Gly	Asp	Gly	Thr	Thr	Thr	Ala	Thr	Val	Leu	Ala
				85				90						95	
Glu	Ala	Ile	Tyr	Ser	Glu	Gly	Leu	Arg	Asn	Val	Thr	Ala	Gly	Ala	Asn
			100					105					110		
Pro	Met	Asp	Leu	Lys	Arg	Gly	Ile	Asp	Lys	Ala	Val	Lys	Val	Val	Val
		115					120					125			
Asp	Glu	Leu	Lys	Lys	Ile	Ser	Lys	Pro	Val	Gln	His	His	Lys	Glu	Ile
	130					135					140				
Ala	Gln	Val	Ala	Thr	Ile	Ser	Ala	Asn	Asn	Asp	Ser	Glu	Ile	Gly	Asn
145					150					155					160
Leu	Ile	Ala	Glu	Ala	Met	Glu	Lys	Val	Gly	Lys	Asn	Gly	Ser	Ile	Thr
				165					170					175	
Val	Glu	Glu	Ala	Lys	Gly	Phe	Glu	Thr	Val	Leu	Asp	Val	Val	Glu	Gly
			180					185					190		
Met	Asn	Phe	Asn	Arg	Gly	Tyr	Leu	Ser	Ser	Tyr	Phe	Ser	Thr	Asn	Pro
		195					200					205			
Glu	Thr	Gln	Glu	Cys	Val	Leu	Glu	Asp	Ala	Leu	Ile	Leu	Ile	Tyr	Asp
	210					215					220				
Lys	Lys	Ile	Ser	Gly	Ile	Lys	Asp	Phe	Leu	Pro	Val	Leu	Gln	Gln	Val
225					230					235					240
Ala	Glu	Ser	Gly	Arg	Pro	Leu	Leu	Ile	Ile	Ala	Glu	Glu	Ile	Glu	Gly
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Glu	Ala	Leu	Ala	Thr	Leu	Val	Val	Asn	Arg	Leu	Arg	Ala	Gly	Phe	Arg
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Val	Cys	Ala	Val	Lys	Ala	Pro	Gly	Phe	Gly	Asp	Arg	Arg	Lys	Ala	Met
	275						280					285			
Leu	Glu	Asp	Ile	Ala	Ile	Leu	Thr	Gly	Gly	Gln	Leu	Val	Ser	Glu	Glu
	290					295					300				
Leu	Gly	Met	Lys	Leu	Glu	Asn	Thr	Thr	Leu	Ala	Met	Leu	Gly	Lys	Ala
305					310					315					320
Lys	Lys	Val	Ile	Val	Thr	Lys	Glu	Asp	Thr	Thr	Ile	Val	Glu	Gly	Leu
				325					330					335	
Gly	Asn	Lys	Pro	Asp	Ile	Gln	Ala	Arg	Cys	Asp	Asn	Ile	Lys	Lys	Gln
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Ile	Glu	Asp	Ser	Thr	Ser	Asp	Tyr	Asp	Lys	Glu	Lys	Leu	Gln	Glu	Arg
	355						360					365			
Leu	Ala	Lys	Leu	Ser	Gly	Gly	Val	Ala	Val	Ile	Arg	Val	Gly	Ala	Ala
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Thr	Glu	Ile	Glu	Met	Lys	Glu	Lys	Lys	Asp	Arg	Val	Asp	Asp	Ala	Gln
385					390					395					400
His	Ala	Thr	Ile	Ala	Ala	Val	Glu	Glu	Gly	Ile	Leu	Pro	Gly	Gly	Gly
				405					410					415	
Thr	Ala	Leu	Val	Arg	Cys	Ile	Pro	Thr	Leu	Glu	Ala	Phe	Leu	Pro	Met
			420					425					430		
Leu	Ala	Asn	Glu	Asp	Glu	Ala	Ile	Gly	Thr	Arg	Ile	Ile	Leu	Lys	Ala
		435					440					445			
Leu	Thr	Ala	Pro	Leu	Lys	Gln	Ile	Ala	Ser	Asn	Ala	Gly	Lys	Glu	Gly
	450					455					460				
Ala	Ile	Ile	Cys	Gln	Gln	Val	Leu	Ala	Arg	Ser	Ala	Asn	Glu	Gly	Tyr
465					470					475					480

Asp	Ala	Leu	Arg	Asp	Ala	Tyr	Thr	Asp	Met	Ile	Asp	Ala	Gly	Ile	Leu
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Asp	Pro	Thr	Lys	Val	Thr	Arg	Ser	Ala	Leu	Glu	Ser	Ala	Ala	Ser	Ile
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Ala	Gly	Leu	Leu	Leu	Thr	Thr	Glu	Ala	Leu	Ile	Ala	Asp	Ile	Pro	Glu
		515					520					525			
Glu	Lys	Ser	Ser	Ser	Ala	Pro	Ala	Met	Pro	Ser	Ala	Gly	Met	Asp	Tyr
	530					535					540				